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Amino acid and protein incorporation in water-stressed perennial ryegrass

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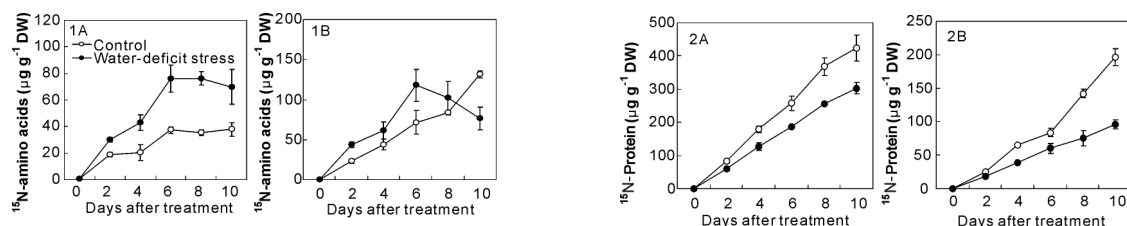
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Introduction During prolonged periods of drought, the decrease in water availability for transport-associated processes leads to changes in the concentrations of many metabolites, followed by disturbances in amino acid and carbohydrate metabolism. In particular, changes in the concentration and composition of the N-soluble fraction in response to both water-deficit (Girousse *et al.* 1996; Lazcano-Ferrat and Lovatt 1999) and salinity stress (Gilbert *et al.* 1998; Silveira *et al.* 2001) have been demonstrated in a wide range of species. Although there have been many studies of the physiological and molecular processes that enable the plant to tolerate drought stress and of nutrient acquisition during water deficit, little information is available concerning the kinetics of *de novo* amino acid and protein synthesis in relation to increasing stress intensity over time, during water deficit. The present work was designed to investigate the kinetic patterns of *de novo* protein and amino acid synthesis in response to increasing intensity of water deficit stress.

Materials and methods ¹⁵NO₃⁻ feeding was carried out daily throughout a 10 day experimental period. For ¹⁵N feeding for the well-watered (control) treatment, 25 mL of ¹⁵N solution (1 mM K¹⁵NO₃ with 8.34 ¹⁵N atom % excess) was administered evenly through three porous plastic tubes buried vertically to a depth of 5 cm in each pot at 10:00 h and 16:00 h, respectively. Using the same protocol of administration, plants submitted to water-deficit received 2.5 mL of ¹⁵N solution, containing the same ¹⁵N atom % and the same amount of N as applied to the control pot (i.e. 0.7 mg N pot⁻¹ d⁻¹). Amino acid and protein were fractionated as described by Kim *et al.* (2004). N content and ¹⁵N atom % of all fractions were determined using N single mode analysis on an ANCA-SL isotopic ratio mass spectrometer (PDZ-Europa, Crewe, UK). Samples were harvested at 0, 2, 4, 6, 8, and 10 days after treatment.

Results In the leaves under water deficit conditions, the amount of newly absorbed N in the amino acid fraction (¹⁵N-amino acid) increased much more rapidly during the initial 6 days and was 2.0-fold higher at day 6 compared to the control. The levels were continuously maintained until day 10 (Figure 1A). In roots, water deficit increased the ¹⁵N-amino acid for the initial 6 days (+65% compared with the control at day 6), and then declined to 58.2% of control values at day 10 (Figure 1B). The amount of N derived from the newly absorbed N in the protein fraction (¹⁵N-protein) in both leaves and roots increased linearly, but the absolute level was significantly lower in water deficit plants. The amount of ¹⁵N-protein in water deficit stressed leaves was 27.2% lower than control after 6 days of water deficit treatment. The ¹⁵N-protein level continued to decrease relative to control until day 10 (71.3%, Figure 2A). *De novo* protein synthesis in roots was significantly lower in water deficit (-51%) than in control plants at day 10 (Figure 2B). These results show that drought stress significantly increased ¹⁵N-amino acid synthesis while it decreased ¹⁵N-protein synthesis. These results suggest that the accumulation of amino acid is possibly due to *de novo* synthesis rather than protein degradation.



Conclusions The present data clearly indicate that the increase in *de novo* amino acid synthesis caused by water deficit is a transient adaptive response for the early period, which is mainly due to the decrease in *de novo* protein synthesis.

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